

<110> Tartaglia, Louis A.

SEQUENCE LISTING

Weng, Xun <120> Nucleic Acid Molecules Encoding GLUTX and Uses Thereof <130> MPI1998-021DV3 <140> US 09/981,947 <141> 2001-10-18 <150> 09/610,417 <151> 2000-07-05 <150> 09/299,549 <151> 1999-04-26 <150> 09/031,392 <151> 1998-02-26 <160> 10 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 2343 <212> DNA <213> Human <220> <221> CDS <222> (73)...(1764) <400> 1 tcqacccacg cqtccqqcct tqqcaqaqtc tqqqqtccct qqactqaqcc atcaqctqqq 60 tcactgagac cc atg qca agg aaa caa aat agg aat tcc aag gaa ctg ggc 111 Met Ala Arg Lys Gln Asn Arg Asn Ser Lys Glu Leu Gly 1 10 cta gtt ccc ctc aca gat gac acc agc cac gcc ggg cct cca ggg cca Leu Val Pro Leu Thr Asp Asp Thr Ser His Ala Gly Pro Pro Gly Pro 15 2.0 ggg agg gca ctg ctg gag tgt gac cac ctg agg agt ggg gtg cca ggt 207 Gly Arg Ala Leu Leu Glu Cys Asp His Leu Arg Ser Gly Val Pro Gly 30 35 45 255 gga agg aga aga aag gac tgg tcc tgc tcg ctc ctc gtg gcc tcc ctc Gly Arg Arg Arg Lys Asp Trp Ser Cys Ser Leu Leu Val Ala Ser Leu gcg ggc gcc ttc ggc tcc tcc ttc ctc tac ggc tac aac ctg tcg gtg 303

| Ala | Gly | Ala | Phe 65 | _ | Ser | Ser | Phe | Leu 70 | Tyr | Gly | Tyr | Asn | Leu 75 | Ser | Val | |
|-----|-----|-----|-----------|-----|-----|-----|-----|-----------|-------------------|-----|-----|-----|-----------|-----|-----|-----|
| - | | - | | | _ | | | _ | gcc Ala | | | | | | | 351 |
| _ | _ | | | | - | | | - | cca Pro | _ | | _ | | _ | | 399 |
| | | | | | | | | - | atc Ile | | | | | | - | 447 |
| | | | | | | | | | ctt Leu 135 | | | | His | | | 495 |
| - | - | | | | | - | | | gct Ala | - | - | | | | | 543 |
| _ | | _ | _ | | _ | | _ | - | ctc Leu | | | | _ | | | 591 |
| _ | | | _ | | | - | _ | | agt Ser | | | | _ | | | 639 |
| | | | | | | | | | ggc Gly | | | | | | | 687 |
| | | | | | | | | | act Thr 215 | | | | | | | 735 |
| | | | | | | | | | tgg Trp | | | | | | | 783 |
| | | _ | | | _ | _ | _ | _ | ctg Leu | - | | | | | | 831 |
| | | | | | | | | | aag Lys | | | | | | | 879 |
| | | _ | | | _ | | _ | | aaa Lys | - | _ | - | | | | 927 |
| gta | gag | gag | gtc | ctg | gct | gag | agc | cac | gtg | cag | agg | agc | atc | cgc | ctg | 975 |

| • | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-------------------|------------|-----|------|
| | Val | Glu | Glu | Val | Leu 290 | Ala | Glu | Ser | His | Val 295 | Gln | Arg | Ser | Ile | Arg 300 | Leu | |
| | | | | | | | | | | | | | | tgg Trp 315 | | | 1023 |
| | | | | | | | | | | | | | | ggc Gly | | | 1071 |
| . • | _ | | | | | | | _ | | | | | - | ggg Gly | | | 1119 |
| | | | | | | | | | | | | | | atc Ile | | | 1167 |
| | | | | | | | | | | | | | | gga Gly | | | 1215 |
| | | | | | | | | | | | | | | ttt Phe 395 | Gly | | 1263 |
| | | | | | | | | | | | | | | gtc Val | | | 1311 |
| | | | | | | | | | | | | | | tgc Cys | | | 1359 |
| | | | | | | | | | | | | | | cag Gln | | | 1407 |
| | _ | | _ | - | - | | | | _ | | | _ | | tgg Trp | | | 1455 |
| · | | | | | | | | | | | | | | agt Ser 475 | | | 1503 |
| | | | | | | | | | | | | | | ggt Gly | | | 1551 |
| | | | | | | | | | | | | | | tat Tyr | | | 1599 |
| | atc | agc | cag | gca | ttt | tcc | aaa | agg | aac | aaa | gca | tac | cca | cca | gaa | gag | 1647 |

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Ile Ser Gln Ala Phe Ser Lys Arg Asn Lys Ala Tyr Pro Pro Glu Glu
                   515
                                       520
                                                                1695
aaa atc gac tca gct gtc act gat gct cct gct tct tct cct ttc act
Lys Ile Asp Ser Ala Val Thr Asp Ala Pro Ala Ser Ser Pro Phe Thr
act ccg aat aca gcc tgg att caa gct gcc gcc acc acc acc gcc acc
                                                                1743
Thr Pro Asn Thr Ala Trp Ile Gln Ala Ala Ala Thr Thr Ala Thr
            545
                               550
aaa aaa gaa cac cca ttg taa acggtcatgt ggtatttcct caacctggaa
                                                                1794
Lys Lys Glu His Pro Leu *
        560
tgaccttccc ctatcttctt ctcctggaga acaccaagtc atgatgtcag acaagagctt 1854
ggattttgga gacatgggtt tgaattccag tcattcattc ttttattcag caaatattta 1914
acaagtactg acatgtccca tatgttgttt tacccactgg ttatacaatg ggagggagag 1974
agagagagag agagagagag agatgctatt ctaaaagctt gaagtctagg ctgtgcacgg 2034
tggctcacgc ctgtaatccc agcactttgg gaggccgagg tgggtggatc gtgaggtcag 2094
qaqattqaqa ccatcctggc taacatggtq aaactccctc tctactaaaa atacaaaaaa 2154
ttagctgagc atggtggcgg gcgcctgtag tcccagctac ttgggaggct gaggcaggag 2214
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Leu Leu Glu Cys Asp His Leu Arg Ser Gly Val Pro Gly Gly Arg Arg
                           40
Arg Lys Asp Trp Ser Cys Ser Leu Leu Val Ala Ser Leu Ala Gly Ala
                       55
Phe Gly Ser Ser Phe Leu Tyr Gly Tyr Asn Leu Ser Val Val Asn Ala
                   70
                                       75
Pro Thr Pro Tyr Ile Lys Ala Phe Tyr Asn Glu Ser Trp Glu Arg Arg
His Gly Arg Pro Ile Asp Pro Asp Thr Leu Thr Leu Trp Ser Val
           100
                               105
                                                   110
Thr Val Ser Ile Phe Ala Ile Gly Gly Leu Val Gly Thr Leu Ile Val
                           120
Lys Met Ile Gly Lys Val Leu Gly Arg Lys His Thr Leu Leu Ala Asn
                       135
Asn Gly Phe Ala Ile Ser Ala Ala Leu Leu Met Ala Cys Ser Leu Gln
145
                   150
                                                           160
                                       155
Ala Gly Ala Phe Glu Met Leu Ile Val Gly Arg Phe Ile Met Gly Ile
                                   170
Asp Gly Gly Val Ala Leu Ser Val Leu Pro Met Tyr Leu Ser Glu Ile
           180
                               185
                                                   190
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Ser Pro Lys Glu Ile Arg Gly Ser Leu Gly Gln Val Thr Ala Ile Phe
   195
                          200
Ile Cys Ile Gly Val Phe Thr Gly Gln Leu Leu Gly Leu Pro Glu Leu
                      215
                                         220
Leu Gly Lys Glu Ser Thr Trp Pro Tyr Leu Phe Gly Val Ile Val Val
                  230
                                     235
Pro Ala Val Val Gln Leu Leu Ser Leu Pro Phe Leu Pro Asp Ser Pro
               245
                                  250
Arg Tyr Leu Leu Glu Lys His Asn Glu Ala Arg Ala Val Lys Ala
           260
                              265
Phe Gln Thr Phe Leu Gly Lys Ala Asp Val Ser Gln Glu Val Glu Glu
                          280
Val Leu Ala Glu Ser His Val Gln Arg Ser Ile Arg Leu Val Ser Val
                      295
                                         300
Leu Glu Leu Leu Arg Ala Pro Tyr Val Arg Trp Gln Val Val Thr Val
                  310
                                     315
Ile Val Thr Met Ala Cys Tyr Gln Leu Cys Gly Leu Asn Ala Ile Trp
                                  330
              325
Phe Tyr Thr Asn Ser Ile Phe Gly Lys Ala Gly Ile Pro Pro Ala Lys
                             345
Ile Pro Tyr Val Thr Leu Ser Thr Gly Gly Ile Glu Thr Leu Ala Ala
       355
                          360
Val Phe Ser Gly Leu Val Ile Glu His Leu Gly Arg Arg Pro Leu Leu
                      375
                                          380
Ile Gly Gly Phe Gly Leu Met Gly Leu Phe Phe Gly Thr Leu Thr Ile
                  390
                                     395
Thr Leu Thr Leu Gln Asp His Ala Pro Trp Val Pro Tyr Leu Ser Ile
              405
                                  410
Val Gly Ile Leu Ala Ile Ile Ala Ser Phe Cys Ser Gly Pro Gly Gly
          420
                              425
Ile Pro Phe Ile Leu Thr Gly Glu Phe Phe Gln Gln Ser Gln Arg Pro
      435
                          440
                                             445
Ala Ala Phe Ile Ile Ala Gly Thr Val Asn Trp Leu Ser Asn Phe Ala
                      455
Val Gly Leu Leu Phe Pro Phe Ile Gln Lys Ser Leu Asp Thr Tyr Cys
                  470
                                      475
Phe Leu Val Phe Ala Thr Ile Cys Ile Thr Gly Ala Ile Tyr Leu Tyr
              485
                                  490
Phe Val Leu Pro Glu Thr Lys Asn Arg Thr Tyr Ala Glu Ile Ser Gln
          500
                              505
Ala Phe Ser Lys Arg Asn Lys Ala Tyr Pro Pro Glu Glu Lys Ile Asp
                         520
                                             525
Ser Ala Val Thr Asp Ala Pro Ala Ser Ser Pro Phe Thr Thr Pro Asn
        535 540
Thr Ala Trp Ile Gln Ala Ala Ala Thr Thr Ala Thr Lys Lys Glu
545
                 <sub>.</sub> 550
                                      555
His Pro Leu
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<211> 383

<212> PRT

<213> Human

<400> 3

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                               25
Met Tyr Val Gly Glu Val Ser Pro Thr Glu Leu Arg Gly Ala Leu Gly
Thr Leu His Gln Leu Gly Ile Val Val Gly Ile Leu Ile Ala Gln Val
                       55
Phe Gly Leu Asp Ser Ile Met Gly Asn Gln Glu Leu Trp Pro Leu Leu
                                       75
Leu Ser Val Ile Phe Ile Pro Ala Leu Leu Gln Cys Ile Leu Leu Pro
              85
                                   90
Phe Cys Pro Glu Ser Pro Arg Phe Leu Leu Ile Asn Arg Asn Glu Glu
                              105
Asn Arg Ala Lys Ser Val Leu Lys Lys Leu Arg Gly Thr Ala Asp Val
                          120
Thr Arg Asp Leu Gln Glu Met Lys Glu Glu Ser Arg Gln Met Met Arg
                      135
                                          140
Glu Lys Lys Val Thr Ile Leu Glu Leu Phe Arg Ser Ala Ala Tyr Arg
                   150
                                      155
Gln Pro Ile Leu Ile Ala Val Val Leu Gln Leu Ser Gln Gln Leu Ser
               165
                                   170
Gly Ile Asn Ala Val Phe Tyr Tyr Ser Thr Ser Ile Phe Glu Lys Ala
           180
                               185
Gly Val Gln Gln Pro Val Tyr Ala Thr Ile Gly Ser Gly Ile Val Asn
                          200
Thr Ala Phe Thr Val Val Ser Leu Phe Val Val Glu Arg Ala Gly Arg
                      215
                                          220
Arg Thr Leu His Leu Ile Gly Leu Ala Gly Met Ala Gly Cys Ala Val
                  230
                                      235
Leu Met Thr Ile Ala Leu Ala Leu Glu Glu Leu Pro Trp Met Ser
              245
                                  250
Tyr Leu Ser Ile Val Ala Ile Phe Gly Phe Val Ala Phe Phe Glu Val
                               265
Gly Pro Gly Pro Ile Pro Trp Phe Ile Val Ala Glu Leu Phe Ser Gln
       275
                           280
Gly Pro Arg Pro Ala Ala Ile Ala Val Ala Gly Phe Ser Asn Trp Thr
                       295
Ser Asn Phe Ile Val Gly Met Cys Phe Gln Tyr Val Glu Gln Leu Cys
                   310
                                      315
Gly Pro Tyr Val Phe Ile Ile Phe Thr Val Leu Leu Val Leu Phe Phe
               325
                                   330
Ile Phe Thr Tyr Phe Lys Val Pro Glu Thr Lys Gly Arg Thr Phe Asp
                               345
Glu Ile Ala Ser Gly Phe Arg Gln Gly Gly Ala Ser Gln Ser Asp Lys
                           360
Thr Pro Glu Glu Leu Phe His Pro Leu Gly Ala Asp Ser Gln Val
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<211> 534

<212> PRT

<213> Human

<400> 4

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Leu Val Leu Ser Val Phe Thr Ala Val Leu Gly Phe Phe Gln Tyr Gly
                                25
           20
Tyr Ser Leu Gly Val Ile Asn Ala Pro Gln Lys Val Ile Glu Ala His
                           40
Tyr Gly Arg Met Leu Gly Ala Ile Pro Met Val Arg His Ala Thr Asn
Thr Ser Arg Asp Asn Ala Thr Ile Thr Val Thr Ile Pro Gly Thr Glu
                   70
                                       75
Ala Trp Gly Ser Ser Glu Gly Thr Leu Ala Pro Ser Ala Gly Phe Glu
                                   90
Asp Pro Thr Val Ser Pro His Ile Leu Thr Met Tyr Trp Ser Leu Ser
           100
                               105
Val Ser Met Phe Ala Val Gly Gly Met Val Ser Ser Phe Thr Val Gly
       115
                           120
Trp Ile Gly Asp Arg Leu Gly Arg Val Lys Ala Met Leu Val Val Asn
                       135
                                           140
Val Leu Ser Ile Ala Gly Asn Leu Leu Met Gly Leu Ala Lys Met Gly
                   150
                                       155
Pro Ser His Ile Leu Ile Ile Ala Gly Arg Ala Ile Thr Gly Leu Tyr
               165
                                    170
Cys Gly Leu Ser Ser Gly Leu Val Pro Met Tyr Val Ser Glu Val Ser
           180
                               185
Pro Thr Ala Leu Arg Gly Ala Leu Gly Thr Leu His Gln Leu Ala Ile
                           200
Val Thr Gly Ile Leu Ile Ser Gln Val Leu Gly Leu Asp Phe Leu Leu
                                           220
                       215
Gly Asn Asp Glu Leu Trp Pro Leu Leu Gly Leu Ser Gly Val Ala
                   230
                                       235
Ala Leu Leu Gln Phe Phe Leu Leu Leu Cys Pro Glu Ser Pro Arg
                                    250
               245
Tyr Leu Tyr Ile Lys Leu Gly Lys Val Glu Glu Ala Lys Lys Ser Leu
                                265
Lys Arg Leu Arg Gly Asn Cys Asp Pro Met Lys Glu Ile Ala Glu Met
                            280
Glu Lys Glu Lys Gln Glu Ala Ala Ser Glu Lys Arg Val Ser Ile Gly
                                            300
                       295
Gln Leu Phe Ser Ser Ser Lys Tyr Arg Gln Ala Val Ile Val Ala Leu
                   310
                                       315
Met Val Gln Ile Ser Gln Gln Phe Ser Gly Ile Asn Ala Ile Phe Tyr
                                   .330
               325
Tyr Ser Thr Asn Ile Phe Gln Arg Ala Gly Val Gly Gln Pro Val Tyr
           340
                                345
Tyr Ala Thr Ile Gly Val Gly Val Val Asn Thr Val Phe Thr Val Ile
                         360
                                                365
Ser Val Phe Leu Val Glu Lys Ala Gly Arg Arg Ser Leu Phe Leu Ala
                       375
Gly Leu Met Gly Met Leu Ile Ser Ala Val Ala Met Thr Val Gly Leu
                   390
                                        395
Val Leu Leu Ser Gln Phe Ala Trp Met Ser Tyr Val Ser Met Val Ala
                                    410
               405
Ile Phe Leu Phe Val Ile Phe Phe Glu Val Gly Pro Gly Pro Ile Pro
                               425
Trp Phe Ile Val Ala Glu Leu Phe Ser Gln Gly Pro Arg Pro Ala Ala
                           440
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Ile Ala Val Ala Gly Phe Cys Asn Trp Ala Cys Asn Phe Ile Val Gly
                     455
Met Cys Phe Gln Tyr Ile Ala Asp Leu Cys Gly Pro Tyr Val Phe Val
                    470
                                        475
Val Phe Ala Val Leu Leu Val Phe Phe Leu Phe Ala Tyr Leu Lys
                485
                                    490
Val Pro Glu Thr Lys Gly Lys Ser Phe Glu Glu Ile Ala Ala Ala Phe
                                505
Arg Arg Lys Leu Pro Ala Lys Ser Met Thr Glu Leu Glu Asp Leu
        515
                            520
Arg Gly Gly Glu Glu Ala
    530
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Ala Thr Ile Gly Ser Phe Gln Phe Gly Tyr Asn Thr Gly Val Ile Asn
                                25
Ala Pro Glu Ala Ile Ile Lys Asp Phe Leu Asn Tyr Thr Leu Glu Glu
Arg Ser Glu Thr Pro Pro Ser Ser Val Leu Leu Thr Ser Leu Trp Ser
                                            60
                        55
Leu Ser Val Ala Ile Phe Ser Val Gly Gly Met Ile Gly Ser Phe Ser
                    70
Val Gly Leu Phe Val Asn Arg Phe Gly Arg Arg Asn Ser Met Leu Ile
                                    90
               85
·Val Asn Leu Leu Ala Ile Ala Gly Gly Cys Leu Met Gly Phe Cys Lys
            100
                                105
Ile Ala Glu Ser Val Glu Met Leu Ile Leu Gly Arg Leu Ile Ile Gly
                            120
Leu Phe Cys Gly Leu Cys Thr Gly Phe Val Pro Met Tyr Ile Gly Glu
                        135
                                            140
Ile Ser Pro Thr Ala Leu Arg Gly Ala Phe Gly Thr Leu Asn Gln Leu
                   150
                                       155
Gly Ile Val Ile Gly Ile Leu Val Ala Gln Ile Phe Gly Leu Lys Val
                                    170
               165
Ile Leu Gly Thr Glu Asp Leu Trp Pro Leu Leu Gly Phe Thr Ile
            180
                                185
Leu Pro Ala Ile Ile Gln Cys Ala Ala Leu Pro Phe Cys Pro Glu Ser
                            200
                                                205
        195
Pro Arg Phe Leu Leu Ile Asn Arg Lys Glu Glu Lys Ala Lys Glu
                        215
                                            220
Ile Leu Gln Arg Leu Trp Gly Thr Glu Asp Val Ala Gln Asp Ile Gln
                                        235
                    230
Glu Met Lys Asp Glu Ser Met Arg Met Ser Gln Glu Lys Gln Val Thr
                                    250
                245
Val Leu Glu Leu Phe Arg Ala Pro Asn Tyr Arg Gln Pro Ile Ile Ile
                                265
Ser Ile Met Leu Gln Leu Ser Gln Gln Leu Ser Gly Ile Asn Ala Val
        275
                            280
                                                285
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Phe Tyr Tyr Ser Thr Gly Ile Phe Lys Asp Ala Gly Val Gln Glu Pro
                       295
                                           300
Val Tyr Ala Thr Ile Gly Ala Gly Val Val Asn Thr Ile Phe Thr Val
                   310
                                       315
Val Ser Val Phe Leu Val Glu Arg Ala Gly Arg Arg Thr Leu His Leu
               325
                                   330
Ile Gly Leu Gly Gly Met Ala Phe Cys Ser Ile Leu Met Thr Ile Ser
           340
                                345
Leu Leu Lys Asp Asn Tyr Ser Trp Met Ser Phe Ile Cys Ile Gly
                           360
Ala Ile Leu Val Phe Val Ala Phe Phe Glu Ile Gly Pro Gly Pro Ile
                       375
                                           380
Pro Trp Phe Ile Val Ala Glu Leu Phe Gly Gln Gly Pro Arg Pro Ala
                                       395
                   390
Ala Met Ala Val Ala Gly Cys Ser Asn Trp Thr Ser Asn Phe Leu Val
               405
                                   410
Gly Leu Leu Phe Pro Ser Ala Thr Phe Tyr Leu Gly Ala Tyr Val Phe
                               425
Ile Val Phe Thr Val Phe Leu Val Ile Phe Trp Val Phe Thr Phe Phe
                           440
Lys Val Pro Glu Thr Arg Gly Arg Thr Phe Glu Glu Ile Thr Arg Ala
                       455
Phe Glu Gly Gln Val Gln Thr Gly Thr Arg Gly Glu Lys Gly Pro Ile
                   470
                                       475
Met Glu Met Asn Ser Ile Gln Pro Thr Lys Asp Thr Asn Ala
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<213> human

<400> 6

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Gln Leu Ala Ile Val Ile Gly Ile Leu Ile Ala Gln Val Leu Gly Leu
          180
                               185
Glu Ser Leu Leu Gly Thr Ala Ser Leu Trp Pro Leu Leu Gly Leu
                           200
Thr Val Leu Pro Ala Leu Leu Gln Leu Val Leu Leu Pro Phe Cys. Pro
                       215
                                           220
Glu Ser Pro Arg Tyr Leu Tyr Ile Ile Gln Asn Leu Glu Gly Pro Ala
                   230
Arg Lys Ser Leu Lys Arg Leu Thr Gly Trp Ala Asp Val Ser Gly Val
                                   250
               245
Leu Ala Glu Leu Lys Asp Glu Lys Arg Lys Leu Glu Arg Glu Arg Pro
                               265
Leu Ser Leu Leu Gln Leu Leu Gly Ser Arg Thr His Arg Gln Pro Leu
                           280
                                               285
Ile Ile Ala Val Val Leu Gln Leu Ser Gln Gln Leu Ser Gly Ile Asn
                       295
                                           300
Ala Val Phe Tyr Tyr Ser Thr Ser Ile Phe Glu Thr Ala Gly Val Gly
                   310
                                       315
Gln Pro Ala Tyr Ala Thr Ile Gly Ala Gly Val Val Asn Thr Val Phe
                                   330
               325
Thr Leu Val Ser Val Leu Leu Val Glu Arg Ala Gly Arg Arg Thr Leu
           340
                               345
His Leu Leu Gly Leu Ala Gly Met Cys Gly Cys Ala Ile Leu Met Thr
                           360
Val Ala Leu Leu Leu Glu Arg Val Pro Ala Met Ser Tyr Val Ser
                       375
                                           380
Ile Val Ala Ile Phe Gly Phe Val Ala Phe Phe Glu Ile Gly Pro Gly
                   390
                                       395
Pro Ile Pro Trp Phe Ile Val Ala Glu Leu Phe Ser Gln Gly Pro Arg
               405
                                   410
Pro Ala Ala Met Ala Val Ala Gly Phe Ser Asn Trp Thr Ser Asn Phe
           420
                               425
Ile Ile Gly Met Gly Phe Gln Tyr Val Ala Glu Ala Met Gly Pro Tyr
                           440
       435
                                               445 ·
Val Phe Leu Phe Ala Val Leu Leu Gly Phe Phe Ile Phe Thr
                       455
Phe Leu Arg Val Pro Glu Thr Arg Gly Arg Thr Phe Asp Gln Ile Ser
                   470
                                       475
Ala Ala Phe His Arg Thr Pro Ser Leu Leu Glu Glu Val Lys Pro
                                   490
Ser Thr Glu Leu Glu Tyr Leu Gly Pro Asp Glu Asn Asp
                               505
           500
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<210> 7
<211> 500
<212> PRT
<213> human
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<400> 7

 Met
 Glu
 Gln
 Asp
 Gln
 Ser
 Met
 Lys
 Glu
 Gly
 Arg
 Leu
 Thr
 Leu
 Val

 Leu
 Ala
 Leu
 Ala
 Thr
 Leu
 Ile
 Ala
 Ala
 Phe
 Gly
 Ser
 Ser
 Phe
 Gln
 Tyr

 Gly
 Tyr
 Asn
 Val
 Ala
 Ala
 Val
 Asn
 Ser
 Pro
 Ala
 Leu
 Leu
 Met
 Gln
 Gln

 35
 Image: Ser
 Image:

```
Phe Tyr Asn Glu Thr Tyr Tyr Gly Arg Thr Gly Glu Phe Met Glu Asp
                       5.5
Phe Pro Leu Thr Leu Leu Trp Ser Val Thr Val Ser Met Phe Pro Phe
                   70
                                       75
Gly Gly Phe Ile Gly Ser Leu Leu Val Gly Pro Leu Val Asn Lys Phe
                                    90
Gly Arg Lys Gly Ala Leu Leu Phe Asn Asn Ile Phe Ser Ile Val Pro
            100
                               105
Ala Ile Leu Met Gly Cys Ser Arg Val Ala Thr Ser Phe Glu Leu Ile
                           120
Ile Ile Ser Arg Leu Leu Val Gly Ile Cys Ala Gly Val Ser Ser Asn
                       135
                                           140
Val Val Pro Met Tyr Leu Gly Glu Leu Ala Pro Lys Asn Leu Arg Gly
                                       155
                   150
Ala Leu Gly Val Val Pro Gln Leu Phe Ile Thr Val Gly Ile Leu Val
                                   170
               165
Ala Gln Ile Phe Gly Leu Arg Asn Leu Leu Ala Asn Val Asp Gly Trp
                               185
Pro Ile Leu Leu Gly Leu Thr Gly Val Pro Ala Ala Leu Gln Leu Leu
                            200
Leu Leu Pro Phe Pro Glu Ser Pro Arg Tyr Leu Leu Ile Gln Lys
                       215
                                            220
Lys Asp Glu Ala Ala Ala Lys Lys Ala Leu Gln Thr Leu Arg Gly Trp
                   230
                                       235
Asp Ser Val Asp Arg Glu Val Ala Glu Ile Arg Gln Glu Asp Glu Ala
                                    250
               245
Glu Lys Ala Ala Gly Phe Ile Ser Val Leu Lys Leu Phe Arg Met Arg
                               265
Ser Leu Arg Trp Gln Leu Leu Ser Ile Ile Val Leu Met Gly Gly Gln
                           280
                                                285
Gln Leu Ser Gly Val Asn Ala Ile Tyr Tyr Tyr Ala Asp Gln Ile Tyr
                       295
Leu Ser Ala Gly Val Pro Glu Glu His Val Gln Tyr Val Thr Ala Gly
                                        315
                    310
Thr Gly Ala Val Asn Val Val Met Thr Phe Cys Ala Val Phe Val Val
                                    330
                325
Glu Leu Leu Gly Arg Arg Leu Leu Leu Leu Gly Phe Ser Ile Cys
           340
                               345
Leu Ile Ala Cys Cys Val Leu Thr Ala Ala Leu Ala Leu Gln Asp Thr
                           360
                                                365
Val Ser Trp Met Pro Tyr Ile Ser Ile Val Cys Val Ile Ser Tyr Val
                        375
                                            380
Ile Gly His Ala Leu Gly Pro Ser Pro Ile Pro Ala Leu Leu Ile Thr
                    390
                                        395
Ile Phe Leu Gln Ser Ser Arg Pro Ser Ala Phe Met Val Gly Gly Ser
                                    410
Val His Trp Leu Ser Asn Phe Thr Val Gly Leu Ile Phe Pro Phe Ile
                                425
            420
Gln Glu Gly Leu Gly Pro Tyr Ser Phe Ile Val Phe Ala Val Ile Cys
                            440
Leu Ile Thr Thr Ile Tyr Ile Phe Leu Ile Val Pro Glu Thr Lys Ala
                       455
                                            460
Lys Thr Phe Ile Glu Ile Asn Gln Ile Phe Thr Lys Met Asn Lys Val
                   470
                                       475
Ser Glu Val Tyr Pro Glu Lys Glu Glu Leu Lys Glu Leu Pro Pro Val
                485
                                    490
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<213> Artificial Sequence
<220>
<223> primer sequence
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                                                                   21
<210> 9
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> primer sequence
<400> 9
ttgttaaggc cttccatt
                                                                   18
<210> 10
<211> 493
<212> PRT
<213> Artificial Sequence
<220>
<223> majority alignment sequence
<221> VARIANT
<222> (1)...(493)
<223> Xaa = Any Amino Acid
<400> 10
Met Xaa Xaa Gly Phe Gln Xaa Gly Ser Val Thr Gly Thr Leu Val Leu
                                    10
Ala Val Leu Ile Ala Ala Leu Gly Ser Phe Gln Tyr Gly Tyr Asn Leu
                                25
Gly Val Ile Asn Ala Pro Gln Lys Val Ile Glu Ala Phe Tyr Glu Thr
                            40
Trp Leu Gly Arg Xaa Gly Glu Xaa Pro Ser Val Pro Thr Leu Thr Leu
                        55
Leu Trp Ser Leu Ser Val Ser Ile Phe Ala Val Gly Gly Met Ile Gly
Ser Phe Leu Val. Gly Xaa Ile Gly Asn Arg Leu Gly Arg Lys Xaa Ala
                85
Met Leu Val Asn Asn Val Leu Ala Ile Ala Gly Gly Leu Leu Met Gly
                                105
Leu Ala Lys Xaa Ala Xaa Ser Phe Glu Met Leu Ile Leu Gly Arg Phe
                            120
```

Thr Ser Glu Gln

Ile Ile Gly Leu Tyr Cys Gly Leu Ser Ser Gly Val Val Pro Met Tyr

```
135
Val Gly Glu Ile Ser Pro Thr Ala Leu Arg Gly Ala Leu Gly Thr Leu
        150
                           155
Asn Gln Leu Gly Ile Val Ile Gly Ile Leu Ile Ala Gln Val Leu Gly
               165
                                  170
Leu Asp Ser Leu Leu Gly Asn Glu Ser Leu Trp Pro Leu Leu Gly
                               185
Leu Thr Gly Val Pro Ala Leu Leu Gln Leu Leu Leu Pro Phe Cys
                          200
Pro Glu Ser Pro Arg Tyr Leu Leu Ile Asn Lys Asn Glu Glu Ala Arg
                      215
Ala Lys Lys Ala Leu Gln Arg Leu Arg Gly Thr Ala Asp Val Ser Gln
                                      235
                   230
Glu Val Ala Glu Met Lys Asp Glu Ser Arg Xaa Met Xaa Ser Glu Lys
              245
                                  250
Xaa Val Ser Val Leu Glu Leu Phe Arg Ser Arg Xaa Tyr Arg Gln Pro
                              265
Val Ile Ile Ala Ile Val Leu Gln Leu Ser Gln Gln Leu Ser Gly Ile
       275
                           280
Asn Ala Val Phe Tyr Tyr Ser Thr Ser Ile Phe Glu Lys Ala Gly Val
                       295
                                          300
Gly Gln Pro Val Tyr Ala Thr Ile Gly Ala Gly Val Val Asn Thr Val
                  310
                                      315
Phe Thr Val Val Ser Val Phe Val Val Glu Arg Ala Gly Arg Arg Thr
                                  330
              325
Leu His Leu Cly Leu Gly Gly Met Ala Gly Cys Ala Val Leu Met
                              345
                                                  350
           340
Thr Ile Ala Leu Ala Leu Leu Asp Gln Val Pro Trp Met Ser Tyr Val
       355
                          360
                                              365
Ser Ile Val Ala Ile Phe Gly Phe Val Ala Phe Phe Glu Val Gly Pro
                       375
                                          380
Gly Pro Ile Pro Trp Phe Ile Val Ala Glu Leu Phe Ser Gln Gly Pro
                   390
                                       395
Arg Pro Ala Ala Ile Ala Val Ala Gly Phe Ser Asn Trp Thr Ser Asn
               405
                                   410
Phe Ile Val Gly Leu Leu Phe Gln Tyr Ile Ala Glu Leu Leu Gly Pro
           420
                              425
                                                  430
Tyr Val Phe Ile Val Phe Ala Val Leu Leu Leu Phe Phe Ile Phe
                          440 ·
Thr Phe Leu Lys Val Pro Glu Thr Lys Gly Arg Thr Phe Asp Glu Ile
                      455
                                          460
Ala Ala Phe Arg Lys Xaa Asn Lys Xaa Glu Gln Pro Glu Lys Glu
                  470
                                      475
Ser Ile Glu Glu Leu Glu Pro Leu Gly Pro Asp Glu Xaa
               485
                                  490
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